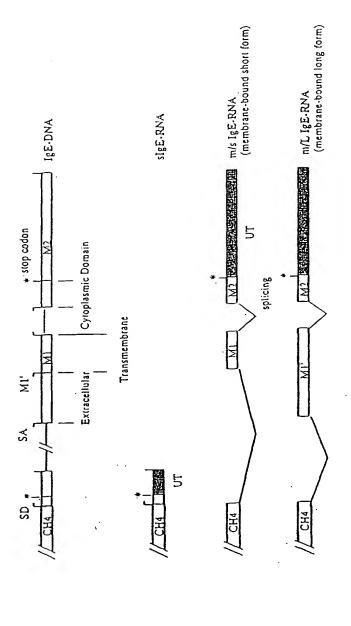
Construction of membrane-bound IgE

I. Cloning of human membrane-bound IgE gene by RT-PCR

A. Schematic representation of the secretary and two membrane human IgE H chain isoforms



B. RT.PCR strategy to amplify m/L IgE form

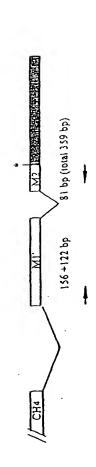
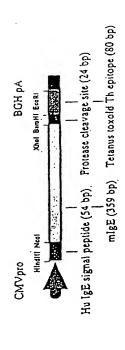


Fig.1

Construction of membrane-bound IgE

II. Construction of a vector insert for expression of a mIgE fused to TT

A. Human mIgE • Tetanus toxoid fusion protein expression cassette



B. PCR amplification with specific primer set

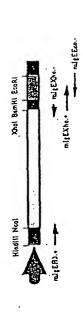


Fig.2

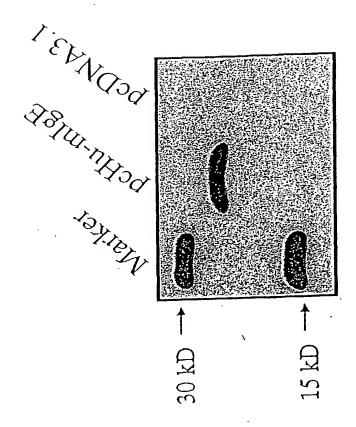


Fig.3

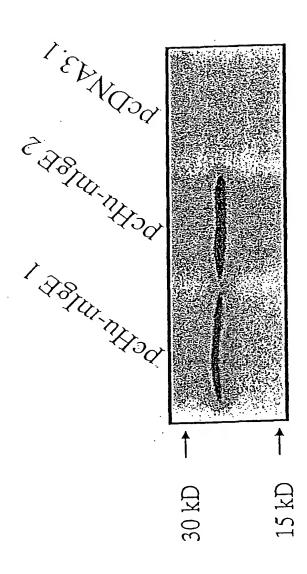
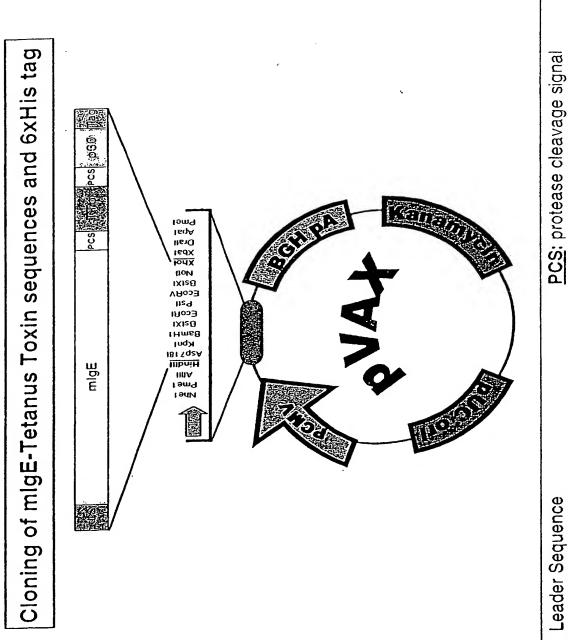


Fig.4



PCS: protease cleavage signal TTp: Tetanus toxin precursor (aa73-99)

7) Tag: 6XHis

migE: human membrane-bound igE p30: Tetanus toxin subunit p30 (aa947-967) . .

10/518701

mlgE-TT construct: Nucleotide and amino acid sequences

(SEQ ID NOS: 5 & 6) atggactggacctggatcctcttcttggtggcagcagccacgcgagtccactcccatggg LFLVAAAT R T Leader sequence ctggctggcggctccgcagtcccagagggccccggatagggtgctctgccactccgga S A Q S Q R A P D R V L C H S G mlaE cagcagcaggagctgccgagagcagcaggaggctctgtcccccacccccgctgccactgt P Н R C H C A G G S Q Q Q GL P R Α ggagccgggagggctgactggccaggtcccccagagctggacgtgtgcgtggaggaggcc L D V C WPGPPE A D GAGR mlgE TGLCIFAALFLL \mathbf{T} \mathbf{W} mlgE agegtgagetacagegeegeeteaegeteeteatggtgeageggtteeteteageeaeg R L S V Q \mathbf{F} \mathbf{T} L M S A A L L mlgE cggcaggggaggccccagacctccctcgactacaccaacgtcctccagccccacgccaga \mathbf{T} S L D Y \mathbf{T} N L Q Р O R O G R migE gaaaaaagagctgttgttggttacgatccaaattatttaaggactgattctgataaagat NY L R G YProtease cleavage signal agatttttacaaaccatggtaaaactgtttaacagaattaagagagaaaaaagagctgtt T M V K L F N R I K R E K R A V Protease cleavage Πρ gttggttttaataattttaccgttagcttttggttgagggttcctaaagtatctgctagt S.F WLRV P K VGFN p30TT signal

catttagaacatcatcatcatcattag

H L E H Flag

Synthetic peptides used as antigens for in vitro assays

Name	Amino acid sequence	Human mlgE Region
IgEEx#1	(SEQ ID NO: 7) SA QSQRAPDR VLCHSGQQQGLP	extracellular (22 aa)
IgEEx#2	(SEQ ID NO: 8) AGGSVPHPRCHCGAGRADWPGP	extracellular (22 aa)
Migis	(SEQ ID NO: 9) ELDVCVEEAEGEAPW	extracellular (15 aa)
CTL2	(SEQ ID NO: 10) EAPWTWTGL	extracellular-transmembrane (9 aa)
כתרו	(SEQ ID NO: 11) TGLCIFAALF	transmembrane (9 aa)
IgECyt	(SEQ ID NO: 12) VQRFLSATRQGRPQTSLDYTNVLQPHA	intracellular (27 aa)
TTh (27 aa) ·	(SEQ ID NO: 13) YDPNYLRTDSDKDRFLQTMVKLFNRIK	

Fig.7

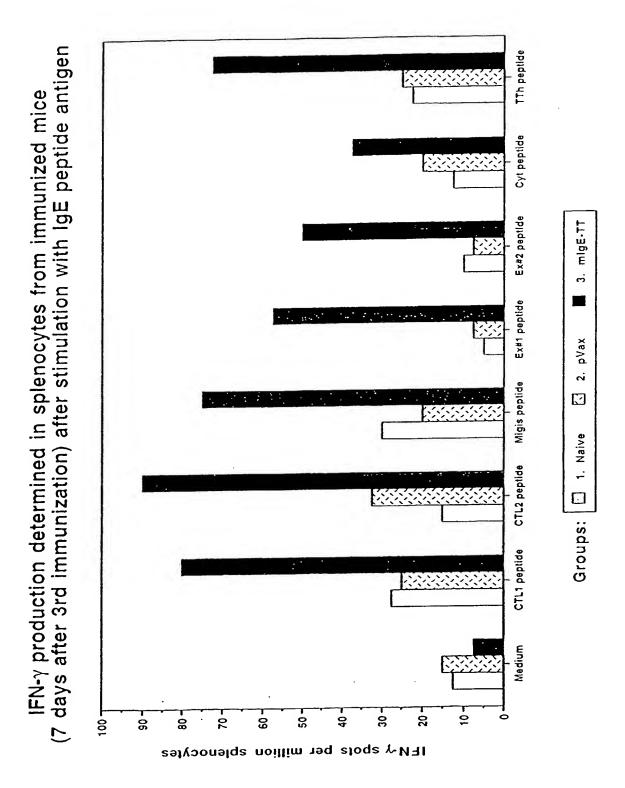


Fig.8